F. Vander Veer

RAW SEQUENCE LISTING DATE: 07/13/2000
PATENT APPLICATION: US/09/323,738 TIME: 15:38:10

Input Set : A:\UW 3570.app
Output Set: N:\CRF3\07132000\I323738.raw

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 3 <110> APPLICANT: Osborne, William R.A.
 4       Ramesh, Nagarajan
 6 <120> TITLE OF INVENTION: Compositions and Methods for Treating Diabetes
 8 <130> FILE REFERENCE: P-UW 3570
10 <140> CURRENT APPLICATION NUMBER: 09/323,738
11 <141> CURRENT FILING DATE: 1999-06-01
13 <150> PRIOR APPLICATION NUMBER: 60/087,660
14 <151> PRIOR FILING DATE: 1998-06-02
16 <150> PRIOR APPLICATION NUMBER: 09/185,852
17 <151> PRIOR FILING DATE: 1998-11-04
19 <160> NUMBER OF SEQ ID NOS: 11
21 <170> SOFTWARE: PatentIn Ver. 2.0
23 <210> SEQ ID NO: 1
24 <211> LENGTH: 450
25 <212> TYPE: DNA
26 <213> ORGANISM: Homo sapiens
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29 <221> NAME/KEY: CDS
30 <222> LOCATION: (45)..(377)
32 <400> SEQUENCE: 1
33 gctgcatcag aagaggccat caagcacatc actgtccttc tgcc atg gcc ctg tgg  56
34                                                 Met Ala Leu Trp
35                                                   1
37 atg cgc ctc ctg ccc ctg ctg gcg ctg ctg gcc ctc tgg gga cct gac  104
38 Met Arg Leu Leu Pro Leu Leu Ala Leu Leu Ala Leu Trp Gly Pro Asp
39   5                  10                  15                  20
41 cca gcc gca gcc ttt gtg aac caa cac ctg tgc ggc tca cac ctg gtg  152
42 Pro Ala Ala Ala Phe Val Asn Gln His Leu Cys Gly Ser His Leu Val
43                  25                  30                  35
45 gaa gct ctc tac cta gtg tgc ggg gaa cga ggc ttc ttc tac aca ccc  200
46 Glu Ala Leu Tyr Leu Val Cys Gly Glu Arg Gly Phe Phe Tyr Thr Pro
47              40                  45                  50
49 aag acc cgc cgg gag gca gag gac ctg cag gtg ggg cag gtg gag ctg  248
50 Lys Thr Arg Arg Glu Ala Glu Asp Leu Gln Val Gly Gln Val Glu Leu
51          55                  60                  65
53 ggc ggg ggc cct ggt gca ggc agc ctg cag ccc ttg gcc ctg gag ggg  296
54 Gly Gly Gly Pro Gly Ala Gly Ser Leu Gln Pro Leu Ala Leu Glu Gly
55      70                  75                  80
57 tcc ctg cag aag cgt ggc att gtg gaa caa tgc tgt acc agc atc tgc  344
58 Ser Leu Gln Lys Arg Gly Ile Val Glu Gln Cys Cys Thr Ser Ile Cys
59  85                  90                  95                  100
61 tcc ctc tac cag ctg gag aac tac tgc aac tag acgcagcccg caggcagccc 397
62 Ser Leu Tyr Gln Leu Glu Asn Tyr Cys Asn
63                  105                 110
65 cccacccgcc gcctcctgca ccgagagaga tggaataaag cccttgaacc agc        450
68 <210> SEQ ID NO: 2
69 <211> LENGTH: 110
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<212> TYPE: PRT
<213> ORGANISM: Homo sapiens
<400> SEQUENCE: 2
Met Ala Leu Trp Met Arg Leu Leu Pro Leu Leu Ala Leu Leu Ala Leu
 1               5                  10                  15
Trp Gly Pro Asp Pro Ala Ala Ala Phe Val Asn Gln His Leu Cys Gly
            20                  25                  30
Ser His Leu Val Glu Ala Leu Tyr Leu Val Cys Gly Glu Arg Gly Phe
        35                  40                  45
Phe Tyr Thr Pro Lys Thr Arg Arg Glu Ala Glu Asp Leu Gln Val Gly
    50                  55                  60
Gln Val Glu Leu Gly Gly Gly Pro Gly Ala Gly Ser Leu Gln Pro Leu
65                  70                  75                  80
Ala Leu Glu Gly Ser Leu Gln Lys Arg Gly Ile Val Glu Gln Cys Cys
                85                  90                  95
Thr Ser Ile Cys Ser Leu Tyr Gln Leu Glu Asn Tyr Cys Asn
            100                 105                 110
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<220> FEATURE:
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<222> LOCATION: (1)..(63)
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Gly Ile Val Glu Gln Cys Cys Thr Ser Ile Cys Ser Leu Tyr Gln Leu
 1               5                  10                  15
gag aac tac tgc aac                                              63
Glu Asn Tyr Cys Asn
            20
<210> SEQ ID NO: 4
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<212> TYPE: PRT
<213> ORGANISM: Homo sapiens
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Gly Ile Val Glu Gln Cys Cys Thr Ser Ile Cys Ser Leu Tyr Gln Leu
 1               5                  10                  15
Glu Asn Tyr Cys Asn
            20
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<212> TYPE: DNA
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ttt gtg aac caa cac ctg tgc ggc tca cac ctg gtg gaa gct ctc tac   48
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```
    139 Phe Val Asn Gln His Leu Cys Gly Ser His Leu Val Glu Ala Leu Tyr
    140  1               5                  10                  15
    142 cta gtg tgc ggg gaa cga ggc ttc ttc tac aca ccc aag acc           90
    143 Leu Val Cys Gly Glu Arg Gly Phe Phe Tyr Thr Pro Lys Thr
    144             20                  25                  30
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    148 <211> LENGTH: 30
    149 <212> TYPE: PRT
    150 <213> ORGANISM: Homo sapiens
    152 <400> SEQUENCE: 6
    153 Phe Val Asn Gln His Leu Cys Gly Ser His Leu Val Glu Ala Leu Tyr
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    157             20                  25                  30
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    162 <212> TYPE: PRT
    163 <213> ORGANISM: Artificial Sequence
    165 <220> FEATURE:
    166 <221> NAME/KEY: SITE
    167 <222> LOCATION: (2)
    168 <223> OTHER INFORMATION: any amino acid
    170 <220> FEATURE:
    171 <221> NAME/KEY: SITE
    172 <222> LOCATION: (3)
    173 <223> OTHER INFORMATION: lysine or arginine or any amino acid (Lys/Arg/Xaa)
    175 <220> FEATURE:
    176 <223> OTHER INFORMATION: Description of Artificial Sequence: Consensus
        177       Sequence
    179 <400> SEQUENCE: 7
W--> 180 Arg Xaa Xaa Arg
    181  1
    184 <210> SEQ ID NO: 8
    185 <211> LENGTH: 4
    186 <212> TYPE: PRT
    187 <213> ORGANISM: Artificial Sequence
    189 <220> FEATURE:
    190 <221> NAME/KEY: SITE
    191 <222> LOCATION: (2)
    192 <223> OTHER INFORMATION: any amino acid
    194 <220> FEATURE:
    195 <221> NAME/KEY: SITE
    196 <222> LOCATION: (3)
    197 <223> OTHER INFORMATION: Lysine or Arginine (Lys/Arg)
    199 <220> FEATURE:
    200 <223> OTHER INFORMATION: Description of Artificial Sequence: Consensus
        201       Sequence
    203 <400> SEQUENCE: 8
W--> 204 Arg Xaa Xaa Arg
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Input Set : A:\Uw 3570.app
Output Set: N:\CRF3\07132000\I323738.raw

205 1
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209 <211> LENGTH: 5
210 <212> TYPE: PRT
211 <213> ORGANISM: Artificial Sequence
213 <220> FEATURE:
214 <223> OTHER INFORMATION: Description of Artificial Sequence: Consensus
215 Sequence
217 <400> SEQUENCE: 9
218 Asp Asp Asp Asp Lys
219 1 5
223 <210> SEQ ID NO: 10
224 <211> LENGTH: 4
225 <212> TYPE: PRT
226 <213> ORGANISM: Artificial Sequence
228 <220> FEATURE:
229 <223> OTHER INFORMATION: Description of Artificial Sequence: Consensus
230 Sequence
232 <400> SEQUENCE: 10
233 Ile Glu Gly Arg
234 1
238 <210> SEQ ID NO: 11
239 <211> LENGTH: 4
240 <212> TYPE: PRT
241 <213> ORGANISM: Artificial Sequence
243 <220> FEATURE:
244 <221> NAME/KEY: SITE
245 <222> LOCATION: (2)
246 <223> OTHER INFORMATION: any amino acid
248 <220> FEATURE:
249 <223> OTHER INFORMATION: Description of Artificial Sequence: Consensus
250 Sequence
252 <400> SEQUENCE: 11
W--> 253 Arg Xaa Lys Arg
254 1

VERIFICATION SUMMARY
PATENT APPLICATION: US/09/323,738

DATE: 07/13/2000
TIME: 15:38:11

Input Set : A:\Uw 3570.app
Output Set: N:\CRF3\07132000\I323738.raw

L:180 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:7
L:204 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8
L:253 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:11